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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total store distribution.

SUMMARIES

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KA Gloric A., Gabrielian A.E., Sea and T.J., Hernandov J.R., Henck J.,

KA Hostin D., Houston K.A., Beal and T.J., Hernandov J.R., Henck J.,

KA Klamel K.E., Erdfrit C.E., Fritt G., Kravitz S., Kuip D., Lai Z.,

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Ka Klamel K.E., Selffrit C.E., Karper C.B., Er Z., Kwatism J.A., Keethom D.,

KA Klamel K.E., Selffrit C.E.
Unery Match
Best Local Similarity
7: Campet
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                                                                                                                                                                                                                                                                      PROSUTE: PS00963; KINOSOMAL
                                                                                                                                                                                                                                                                                                                                                           Priam; Probais; Ribosomal, S. PRINES; PROCES; RIBOSOMALS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   EMBL: AE004421; AAF45638.1:
FlyBase; FBqn0003517; sta.
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NCRE_Tax1D 7227;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  01-MAY-2000 (TEEMBLE OL. 18.
01-MAY-2000 (TEEMBLE OL. 18.
01-JUN-2001 (TEEMBLE OL. 17.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MEDILINE 20196006, Foldword (1771142)
Adduns M.D., Celoiker S.E., Belt R.A., Evans C.A., Gacayne J.D.,
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01 - YAY - 10
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                                                                                                                                                                                                                                                                                                                                                                                                                                               InterPro: IPRO01865; Ribosomai S2.
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                                                                                                                                                                                                                                   (7.74.7 MW)
                                                                                                                                                                                                                                                                      Last sequence update)
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                                                                                                                                                                                                                                   2B76750836138B64 ('RC'64;
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COMME 171, 170

Suld*: Store 41; DB 70.0%; Faed. No. 29; Flor D. Mirmatrhea

DB 5: Length (30)

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1.82416
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01-NoV-1958 (Trimmirel, 08, Greated)
01-JUN-2001 (Trimmirel, 17, 15st annotation update)
TYROSINE/1-DA HETAHROXYLASE (FC 4 1_1 15s).
                                                                                                                                                                                                                                                                                                                                        tobacco.
Flant Ph
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InterPro: IPR03456: AAA_Sub-
InterPro: IPR03496: AAA_Sub-
InterPro: IPR03496: AAA_I
Ptam: PF00004: AAA_I
SMART: SM00382: AAA; 1.
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                                                                                                 Mendel: 33066; Parko:1122,34886.
InterPro, IPP001129; PyFidoxal_Med.
Pram: PP00282; PyFidoxal_Med; 1.
PRINTS; PR0085; YHL-MP WLASE.
                                                                                                                                                                                                                                                                                                                                                                                                  MRECINE 98434205, Inchmed 5738527,
Farchiai P.J., Febree Fost C., Samanani N., Kowalchuk B.;
"Papersoles patterns of Served by "paper some road in transaction decarboxylase promothers from option paper are manuscrad in transaction
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Bukaryota: Viriliplante: Streptophyta: Embayophyta: Tracheephyta:
Spermatophyta: Magneliophyta: endicatyledomu. Ranneulales.
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01-0CT-2000 (TERMBLES, 17, Last sequence)
01-0CN-2001 (TERMBLES, 17, Last annotation the content of the c
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                                                                                                                                                                                                                                   -1- SIMILABITY: PYRIDOXAL-DEPENDENT DECARDOXYLASE FAMILY EMBL: AF025434; AA-161843.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SMART; SM00382; AAA; 1
PROSITE; PS00674; AAA;
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EMBL; AFF27592; AATGIE46.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Yao Y., Wand C.C
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SEQUENCE 531 AA; 6
                                                             PROSITE: PSO0 692: DDC_GAD_HDC_YDC: 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STRAIN MARIANNE;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE FROM N.A.
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The COPACTOR PREPIOSEAL PROSPEATE (DV SIMILARITY
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"al Similarity Tile%;
"A. Compose ation
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e. Pyridoxal phosphate.
| $4091_MW:= E6FEE6H7A2270056|CEC64;
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15. Last sequence update)
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Hest Local Similarity
From 2: Conser
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Burtis K.T., Busan D.A., Barter H
Cherry J.M., Cawley D., Gracke C.,
de Pables B., Delmer A., State C.,
Bodson K., Boup L.E., Bownes M. F
Durbin K.J., Evansellista C.C., Ferro
                                                                                                  "The geneme sequence of its signification e. 287:2185, 2195 (2000), EMBG, AEG, 0722 (2000), AMP (587.1)
                                                                                                                                                                                 Stinshas and Technology General Action (Action Market Mark
                                                                                                                                                                                                                                                                                                                   Shue B.C., Siden Klamas I.,
Spier E., Spradling A.C.,
Stilphas at Testar T. in
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Wan K.H.
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Adams M.D., Ugelruker S.E., Berrie H.
Amandalides P.O., retherer S.E., E.
George R.A., Lewis S.F., Returns
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01-MAY-2000 (TEMBELTAL 16 Cost sequence 16.
01-JUN 2001 (FEMBELTAL 17 Lost sequence 16.
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NCBL_laxII 7227;
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Brosophila melan-master (Frd? t.v
REGULEROLD
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HTT., ITRGGIST?.
5 PFU1703; VMATE
HRCE 532 AA **
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Liu S., Li J., Araujo R., Au M., Brendel V., Buchler E., Comway A.B
Conway A.B., Jowar K., Pest J. Kim C., Kurtz D., Li Y., Palm C.J.,
Shimo F., San B., Davis E.W., Ecker J.R., Federspiel N.A.,
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Brinchan R.C., Sykes S.M., Mache T.M., Ketlavade A.R., Adams M.D.,
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